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On the fixation or nonfixation of inversions under epistatic selection

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Abstract

Several recent publications have stated that epistatic fitness interactions cause the fixation of inversions that suppress recombination among the loci involved. Under this type of selection, however, the suppression of recombination in an inversion heterozygote can create a form of heterozygote advantage, which prevents the inversion from becoming fixed by selection. This process has been explicitly modelled by previous workers.

KEYWORDS

balanced polymorphism, epistasis, fitness, inversions, linkage disequilibrium recombination suppression

There is a growing interest in the evolutionary role of inversions and other types of chromosome rearrangements, with several recent review papers (e.g., Faria et al., 2019; Huang & Rieseberg, 2020; Kapun & Flatt, 2019; Wellenreuther & Bernatchez, 2018) and a special issue of *Molecular Ecology* devoted to this topic (Wellenreuther et al., 2019, and references therein). As these papers have pointed out, there are multiple ways in which natural selection can act on inversions. In order to discriminate between different hypotheses, it is necessary to have a clear understanding of their observable consequences. Unfortunately, there appears to be a serious misconception about one process that has been proposed as providing a selective advantage to an inversion. This involves Dobzhansky's (e.g., Dobzhansky, 1949, 1950, 1951) concept of "coadaptation" among polymorphic loci that interact in their effects on fitness, an idea that traces back to Fisher (1930, pp. 102–104).

In its simplest form, this model invokes two loci, A and B, each segregating for a pair of alleles (A_1 vs. A_2 , B_1 vs. B_2) in a diploid, randomly mating population. Assume that the fitness effects of the two loci are epistatic, in the sense that the fitnesses of the nine possible diploid genotypes at the two loci deviate from those predicted by additive combinations of the effects of the alleles at each locus

(Fisher, 1918). It is then possible for linkage disequilibrium (LD) to be maintained at a stable equilibrium in the face of recombination, where both loci are polymorphic, and fitter combinations of alleles are in excess of the frequencies expected by randomly combining alleles according to their frequencies (Fisher, 1930; Karlin, 1975; Kimura, 1956; Lewontin & Kojima, 1960). There is then selection for modifiers that reduce the rate of recombination between the two loci (Feldman, 1972; Fisher, 1930; Kimura, 1956). The same principle applies to more general multilocus systems in randomly mating populations (Charlesworth, 1976; Zhivotovsky et al., 1994), but not necessarily to partially inbreeding populations (Charlesworth et al., 1979). In particular, an inversion that arises on a haplotype that is present in excess of random expectation (and is thus fitter than average) experiences a selective advantage if recombination is suppressed in heterozygotes for the inversion, simply because the inversion maintains its association with a high-fitness genotype (Kimura, 1956; Charlesworth & Charlesworth, 1973).

The misconception is that this process causes the inversion to spread to fixation, implying that it cannot explain the balanced inversion polymorphisms that have been the subject of so much recent attention. This claim appears to have originated in table 1 of

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Kirkpatrick and Barton (2006), and has been repeated in review papers by Hoffmann and Rieseberg (2008), Kapun and Flatt (2019) and Huang and Rieseberg (2020). However, it overlooks the fact that crossing over is suppressed only in inversion heterozygotes, so that the selective benefits of recombination suppression confer a heterozygote advantage to the inversion, and hence prevent its fixation. This scenario was first modelled by Kimura (1956) and examined in more detail by Charlesworth (1974), who showed that the exchange of alleles among inverted and standard arrangements by gene conversion or double crossing over does not prevent the establishment and maintenance of an inversion polymorphism when there is epistatic selection. An example of this mechanism for maintaining inversion polymorphisms is provided by the epistatic interactions among the different components of segregation distorter systems, and probably explains the frequent association of such systems with inversions (Fuller et al., 2020). Direct effects of inversions in causing segregation distortion should, therefore, not be assumed without further evidence. Of course, there are processes other than epistatic selection that can select for inversions, as described in the reviews cited above, and we do not mean to imply that epistatic selection is necessarily involved in the maintenance of inversion polymorphisms.

AUTHOR CONTRIBUTIONS

Both authors wrote the paper.

DATA AVAILABILITY STATEMENT

No new data were generated in this study.

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